

EXHIBIT

9

No.	Target file	Definition	Match%	Over.	INIT	OPT
1	SmW_H_AA_C		92.6	497	1897	2385
		280 290 300 310 320 330				
SmW_R_AA	GDLGGRKTQQRWTTFLKADLLCPQPEHQRASGVLDAAELRPPQAGTPIFYGIFSSQW					
	X::					
SmW_H_AA_C	GDLGGRKTQQRWTTFLKADLLCPQPEHQRASSVLQDVAVLRPELQAGTPIFYGIFSSQW					
	10 20 30 40 50 60					
	340 350 360 370 380 390					
SmW_R_AA	EGAAISAVCAFRPQDIRAVLNGPFRELKHCNRLPVMDNEVPQPRPGECIANMKLQQF					
	::					
SmW_H_AA_C	EGATISAVCAFRPQDIRTVLNGPFRELKHCNRLPVVDNDVPQPRPGECITNNMKLRHF					
	70 80 90 100 110 120					
	400 410 420 430 440 450					
SmW_R_AA	GSSLSLPDRVLTFRDHPLMDRPVFPADGRPLLVTDTAYLRVVAHRVTSLSQKEYDVL					
	::					
SmW_H_AA_C	GSSLSLPDRVLTFRDHPLMDRPVFPADGHPLLVTDTAYLRVVAHRVTSLSQKEYDVL					
	130 140 150 160 170 180					
	460 470 480 490 500 510					
SmW_R_AA	LGTEGHLHRAVRIGAGLSVLEDLALFPEPQPVESMKLYHDMLLVGSHTVTVQVNTSNG					
	::					
SmW_H_AA_C	LGTEGHLHRAVRIGAGLSVLEDLALFPEPQPVENMKLYHSMLLVQSRTEVTQVNTTNG					
	190 200 210 220 230 240					
	520 530 540 550 560 570					
SmW_R_AA	RLQSCSECI LAQDPVCAWFRLDACVAHAGEHRGMVQDIESADVSSLCPEKPEHPVVFE					
	::					
SmW_H_AA_C	RLQSCSECI LAQDPVCAWFRLDACVAHAGEHRLVQDIESADVSSLCPEKPEHPVVFE					
	250 260 270 280 290 300					
	580 590 600 610 620 630					
SmW_R_AA	VPVATVGHVVLPCSPSSAWASCVWHQPSQVLTALTPRRDQLEVVVTPGANGAYACEQEGG					
	::					
SmW_H_AA_C	VPVATAAHVVLPCSPSSAWASCVWHQPSQVLTALTPRRDQLEVVVTPGANGAYACEQEGG					
	310 320 330 340 350 360					
	640 650 660 670 680 690					
SmW_R_AA	AARVVAAYSLVWGSQRGPSNRAHTVVGAGLVGFLGLAASLTLLIGRRQRRRRQRELL					
	::					
SmW_H_AA_C	AAHVVAAYSLVWGSQRDAPSRAHT-VGAGLAGFGLAASLTLLIGRRQRRRRQRELL					

	370	380	390	400	410	
	700	710	720	730	740	750
SmH_R_AA	ARDKVGLDLGAPPSGTTSYSQDPPSPSPEDERLPLALGKRGSGFGGFPPFLLDSCPSA					
	:.....^v:.....					
SmH_H_AA_C	ARDKVGLDLGAPPSGTTSYSQDPPSPSPEDERLPLALAKRGSGFGGFSPFFLLDPCPSA					
	420	430	440	450	460	470
	760	770				
SmH_R_AA	HIRLTQAPLATCETS					
	:.....					
SmH_H_AA_C	HIRLTQAPLATCETS					
	480	490				